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 Trp Cys Ser Ser Trp Leu Tyr Pro Ile His Lys Lys Ser Gly Gly Trp
 40 35 40 45
 Phe Arg Leu Trp Phe Asp Ala Ala Val Leu Leu Ser Pro Phe Thr Arg
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 Cys Phe Ser Asp Arg Met Met Leu Tyr Tyr Asp Pro Asp Leu Asp Asp
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 Tyr Gln Asn Ala Pro Gly Val Gln Thr Arg Val Pro His Phe Gly Ser
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 Thr Lys Ser Leu Leu Tyr Leu Asp Pro Arg Leu Arg Asp Ala Thr Ser
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 Tyr Met Glu His Leu Val Lys Ala Leu Glu Lys Lys Cys Gly Tyr Val
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Ala Ala Ser Gly His Pro Ser Arg Val Ala Ser Gln Phe Leu Gln Asp
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Val Ile Leu Leu Ser His Ser Leu Gly Gly Leu Phe Val Leu His Phe
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10 Leu Asn Arg Thr Thr Pro Ser Trp Arg Arg Lys Tyr Ile Lys His Phe
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Val Ala Leu Ala Ala Pro Trp Gly Gly Thr Ile Ser Gln Met Lys Thr
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20 Val Arg Arg His Gln Arg Thr Ser Glu Ser Asn Gln Trp Leu Leu Pro
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Ser Thr Lys Val Phe His Asp Arg Thr Lys Pro Leu Val Val Thr Pro
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25 Gln Val Asn Tyr Thr Ala Tyr Glu Met Asp Arg Phe Phe Ala Asp Ile
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Gly Phe Ser Gln Gly Val Val Pro Tyr Lys Thr Arg Val Leu Pro Leu
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30 Thr Glu Glu Leu Met Thr Pro Gly Val Pro Val Thr Cys Ile Tyr Gly
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Arg Gly Val Asp Thr Pro Glu Val Leu Met Tyr Gly Lys Gly Phe
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Asp Lys Gln Pro Glu Ile Lys Tyr Gly Asp Gly Asp Gly Thr Val Asn
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40 Leu Ala Ser Leu Ala Ala Leu Lys Val Asp Ser Leu Asn Thr Val Glu
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Val Glu His Met Ser Leu Asp Asn Glu Thr Gly Leu Asp Pro Ala Gly
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Ile Arg Val Arg Ala Val Ser Gly Leu Val Ala Ala Asp Tyr Phe Ala
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10 Pro Gly Tyr Phe Val Trp Ala Val Leu Ile Ala Asn Leu Ala His Ile
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Gly Tyr Glu Glu Lys Asn Met Tyr Met Ala Ala Tyr Asp Trp Arg Leu
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15 Ser Phe Gln Asn Thr Glu Arg Asp Gln Thr Leu Ser Arg Met Lys Ser
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Ala Lys Tyr Ile Lys Ala Val Met Asn Ile Gly Gly Pro Phe Leu Gly
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 210 215 220

Gly Asp Thr Ile Trp Gly Gly Leu Asp Trp Ser Pro Glu Leu Pro Asn
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 245 250 255

Glu Arg Ala Tyr Val Tyr Lys Leu Asn Gln Ser Pro Asp Ser Cys Ile
 260 265 270

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50 Leu Lys Ala Gly Val Tyr Asn Val Asp Gly Asp Glu Thr Val Pro Val
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Leu Ser Ala Gly Tyr Met Cys Ala Lys Ala Trp Arg Gly Lys Thr Arg
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55 Phe Asn Pro Ser Gly Ile Lys Thr Tyr Ile Arg Glu Tyr Asn His Ser
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Pro Pro Ala Asn Leu Leu Glu Gly Arg Gly Thr Gln Ser Gly Ala His
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	Lys Arg Asp Gly Asn Gly Arg Lys Arg Trp Arg Asp Ser Arg Arg Leu			
	65 70 75 80			
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	Gly Ala Tyr His Val His Asn Ser Asp Ser Asp Leu Phe Asp Asn Phe			
	100 105 110			
55	gta aat ttt gat tca ctt aaa gtg tat ttg gat gat tgg aaa gat gtt			384
	Val Asn Phe Asp Ser Leu Lys Val Tyr Leu Asp Asp Trp Lys Asp Val			
	115 120 125			
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	Leu Pro Gln Gly Ile Ser Ser Phe Ile Asp Asp Ile Gln Ala Gly Asn			
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	Tyr Ser Thr Ser Ser Leu Asp Asp Leu Ser Glu Asn Phe Ala Val Gly			
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1986

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Lys Arg Asp Gly Asn Gly Arg Lys Arg Trp Arg Asp Ser Arg Arg Leu
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10 Ile Gly His Ser Met Gly Ser Gln Ile Ile Phe Tyr Phe Met Lys Trp
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55 Val Phe Leu Thr Glu Gly Asp Gly Thr Val Pro Leu Val Ala His Ser
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 Asp Gly Asn Asn Lys Gly Gly Ser Val His Asn Lys Arg Glu Ser Arg
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 Gly Ala Tyr His Val His Asn Ser Asp Ser Asp Leu Phe Asp Asn Phe
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gta aat ttt gat tca ctt aaa gtg tat ttc gat gat tgg aaa gat gtt 384
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ctc cca caa ggt aca agt tcg ttt acc gat gat att cag gct ggt aac 432
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 Gly Ser Phe Tyr Met Leu Arg Thr Met Val Met Asp Lys Val Cys Trp
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1986

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TGTTGTGACT	TACTGGATTG	AGCTCGATAC	CTGATTGTTG	GTTGATTAG	1300
GAGCTCCTCT	TCTGGTTCT	GTGAGGCAA	TCAAATCTAC	TCTCTCTGGT	1350
GTAACGTTTG	GCCTTCCTGT	TTCTGACGTG	ACCTCTGACT	TCTCTTGT	1400
TTTAAGTAGT	TGATATCAAC	CAGGTCTTAT	AACTCACTGG	ATTTCTCTT	1450
TGAAAGTATT	ACTTTGTGA	ATTGAATCTG	TGACGCGAT	ATGGTATCTG	1500
TAGATCTTGA	AGTGTAGTT	ATCAAAGAAC	ATATTGTGGG	TAGTATAACCT	1550
GTCAGCGGCC	TTAGCTAATA	CAACCAAACC	ACATGTACAC	TGATTIAGTT	1600
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CTTITATT	TAATAGGCTA	TGATTGTTT	ATTGAATCA	TGTGACATAT	1700
TGACATGGCG	TTCTCATGTT	TTTGTGTC	AAGGCTTCAG	GGAACTGCTC	1750
GGTGTGTC	CAATTCTTT	GCCTCGTCAT	TGIGGCTTAT	GCCATTTCA	1800
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CAAAATATTC	TGGCTGGCCG	ACAAATATTA	TTAACATTGA	AATTCTTCC	1950
ACTAGCGGTT	AGACTCTGTA	TAIGCAACTG	TAACACTAA	AAAAGTTCA	2000
CCAAGAATGT	TCACCTCTAT	ATTCGTTCC	TTGATGTTG	ATCCATCACT	2050
TACAGAAACA	GCTCTAGTCA	ACATGACCAG	CATGGAATGT	GGCCTTCCCA	2100
CCCTTTGTC	TTTCAGCC	CGTGAACCTAG	CAGATGGGAC	TCTTTCAAA	2150
GCAATAGAAG	ACTATGACCC	AGATAGCAAG	AGGAAGTTAC	ACCAAGTTAA	2200
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ATTCCTAAAT	TCACATTATG	CGTTGACTTT	GTATTATAT	TCCCCATTTG	2700
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CCTATTAGC	GTAAAGGTA	CTAAATGTAT	GAAGCTGTCT	GTCATAGGT	2850
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GTGGCAIGTT	ATCTCAGTTG	CATAAGCAAA	TTATTAACAA	ACTAAAATT	3050
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ACCAAAACAA	AACTARCCC	TTTCTGAA	TCATATTATT	AGGAGTAGTC	3200
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 <211> 709
 <212> cDNA
 <213> tomato
 <400> 11

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CATTCCCAAT	ATGACAAAAGT	TACCTACAT	GAAGTACATA	ACCTATTATG	150
AGGATTCTGA	AAGTTTCCA	GGGACAGAGAA	CAGCAGTTG	GGAGCTTGAT	200
AAAGCAATTC	ACAGGAACAT	TGTCAGACT	CCAGCTTGA	TGCGGGAGCT	250
GTGGCTTGAG	ATGTGGCATG	ATATTG	TGATAAAAAG	TCCAAGTTG	300
TTACAAAAGG	TGGTGTCTGA	TCCTCACTAT	TTTCTTCTAT	AAATGTTGA	350
GTTTGTATTG	ACATTGTAAG	TATTGCAACA	AAAAGCAAAG	CGTGGGCCTC	400
TGAGGGATCA	GGACTGCTAT	TGGGATTACG	GGAAAGCTCG	ATGTGCATGG	450
GCTGAACATT	GTGAATACAG	GTTAGAATAT	TCAATTATA	TTTGCAAAA	500
TATTCTCTTT	TTGTGTATTG	AGGCCACCTT	TCCCCGGTCA	CAACGATGCA	550
GATATGTATT	CGGGGATGTT	CACCTGGGAC	AGAGTTGCAG	ATTGAAGAGT	600
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